

## Blood Group, Immunity, and Risk of Infection with *Vibrio cholerae* in an Area of Endemicity

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**Individuals with blood group O are more susceptible than other individuals to severe cholera, although the mechanism underlying this association is unknown. To assess the respective roles of both intrinsic host factors and adaptive immune responses that might influence susceptibility to infection with *Vibrio cholerae*, we prospectively followed a cohort of household contacts of patients with cholera in Bangladesh. In this study, we made the novel observation that persons with blood group O were less likely than those with other blood groups to become infected with *V. cholerae* O1 (odds ratio [OR], 0.67; 95% confidence interval [CI], 0.53 to 0.85;  $P = 0.008$ ). Consistent with prior studies, however, household contacts with blood group O were more likely to develop severe illness if infected with *V. cholerae* O1 (OR, 2.3; 95% CI, 0.98 to 5.59;  $P = 0.05$ ). While blood group O protected significantly against infection with *V. cholerae* O1, there was no evidence of protection against *V. cholerae* O139. A multivariate analysis demonstrated that the association between blood group O and protection from infection with *V. cholerae* O1 was independent of age, gender, and baseline anti-cholera toxin and vibriocidal antibody titers. Based on this epidemiologic evidence, we propose a hypothesis for understanding the association between blood group O and the risk of infection with *V. cholerae* O1 and O139 as well as the risk of developing severe symptoms once infected.**

*Vibrio cholerae* is a gram-negative bacterium that causes a spectrum of infection ranging from asymptomatic colonization to rapidly fatal secretory diarrhea known as cholera gravis. *V. cholerae* is differentiated serologically by the O side chain of its lipopolysaccharide; the vast majority of human cholera is caused by the O1 and O139 serogroups. The O1 serogroup of *V. cholerae* is subclassified into two biotypes (classical and El Tor) and two major serotypes (Inaba and Ogawa) (11). Due to variations in the predominating serogroup, biotype, and serotype in circulation, the epidemiology of cholera is in constant flux. In the 1960s, the *V. cholerae* O1 El Tor biotype emerged as a major cause of cholera, ultimately replacing the classical biotype. In 1992, the *V. cholerae* O139 serogroup first appeared; after briefly predominating in South Asia, it now persists in this region, but at much lower levels than *V. cholerae* O1 El Tor.

Susceptibility to infection with *V. cholerae* is dependent on both adaptive immune responses induced by previous infection and innate host factors. The best-studied correlate of adaptive immunity to *V. cholerae* is the serum vibriocidal antibody, a complement-fixing bactericidal antibody. Seroepidemiologic studies in areas of endemicity have shown that vibriocidal antibody titers increase with age and that risk of disease is

inversely proportional to the vibriocidal antibody titer (9, 16, 17). However, there is no threshold vibriocidal antibody titer above which complete protection from infection is achieved, and it is hypothesized that the vibriocidal antibody is a surrogate marker for a protective mucosal immune response (24). Systemic antibodies to cholera toxin have not been found to correlate with protection from cholera (9). Furthermore, infection with *V. cholerae* O1 does not confer protection from *V. cholerae* O139, even though both serogroups produce identical cholera toxins (1, 22).

Among the intrinsic host factors that influence susceptibility to cholera, the ABH histo-blood group antigens are the most studied. Multiple case-control studies in areas of cholera endemicity have demonstrated that individuals with blood group O (the blood group phenotype associated with the H antigen) are at increased risk of hospitalization due to classical and El Tor *V. cholerae* O1 as well as *V. cholerae* O139 (3, 4, 5, 7, 26). A study of North American volunteers also demonstrated increased purging in blood group O subjects infected with a high inoculum of classical *V. cholerae* O1 (13). It has been hypothesized that *V. cholerae* infection is the evolutionary force behind the low prevalence of the O blood group in the Ganges River Delta, which is a historic and current global epicenter of cholera (10, 11).

Two previous studies have addressed the question of whether blood group O is also associated with an increased risk of asymptomatic infection or mild illness due to *V. cholerae* in addition to the increased risk of severe illness (10, 26). Both

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studies concluded that there was no difference in the overall risk of infection with *V. cholerae* among persons with blood group O compared to those with non-blood group O antigens. A prospective study of household contacts of cholera patients in Bangladesh performed between 1980 and 1982 found that while blood group O was associated with an increased risk of moderate to severe cholera, there was no association between blood group and the risk of overall infection (defined microbiologically) due to *V. cholerae* O1 El Tor (10). Similar findings were reported in a retrospective evaluation of cholera patients performed during the first outbreak of *V. cholerae* O1 El Tor in Trujillo, Peru (26). In this study, individuals with blood group O did not have increased evidence of recent infection (defined serologically) but were more than eight times as likely to require hospitalization for cholera.

Although the mechanism by which blood group influences susceptibility to cholera is undefined, the association is of significant public health importance. During the epidemic of cholera in the 1990s in South America, where there is a high prevalence of blood group O, the requirements for hospitalization, intravenous fluids, and oral rehydration therapy were greater than seen in outbreaks in other regions (26). The blood group distribution in a population may also influence the efficacy of cholera vaccination programs, although current data are conflicting. A trial of a cholera toxin B subunit-killed oral vaccine in Bangladesh showed a lower protective efficacy in individuals with blood group O (5). In contrast, a live oral attenuated cholera vaccine, CVD 103-HgR, resulted in higher vibriocidal antibody responses in Indonesian individuals with blood group O (12). Although CVD 103-HgR had an overall protective efficacy of only 14% over the 4-year surveillance period, there was a trend toward greater efficacy in individuals with blood group O, in whom the protective efficacy was 46% (23).

To understand the factors that mediate susceptibility to symptomatic and asymptomatic infection with *V. cholerae*, we undertook a prospective, observational study of a cohort of cholera patients and their household contacts in Dhaka, Bangladesh. Here, we present novel findings regarding the association between blood group, the vibriocidal antibody response, and the risk of *V. cholerae* infection in a current area of endemicity.

#### MATERIALS AND METHODS

**Study design and subject enrollment.** The International Centre for Diarrheal Disease Research (ICDDR,B) hospital cares for approximately 10,000 to 20,000 cholera patients annually, the majority of whom are residents of Dhaka city. A random sample of patients over the age of 6 months presenting to the ICDDR,B hospital with severe acute watery diarrhea, dark-field positive stool, and no significant comorbid conditions was selected for inclusion in this study. Within 4 to 6 h of presentation of the index case, a field team discussed enrollment with household contacts of the index patient, defined as individuals sharing the same cooking pot. Blood specimens for ABO typing and baseline vibriocidal and anti-cholera toxin antibody titers were collected immediately upon enrollment from consenting contacts. Contacts were visited by the field team on each of the next 6 days and again on days 14 and 21 after presentation of the index case. During these visits, contacts were questioned about diarrheal symptoms, and rectal swabs were obtained for *V. cholerae* culture. Follow-up blood samples for vibriocidal antibody titers were obtained from contacts on study days 4 and 21. Patients and contacts were excluded from the analysis if they did not complete 21 days of follow-up. Household contacts with diarrhea during the week prior to enrollment or with a positive rectal swab culture for *V. cholerae* upon enrollment

in the study were excluded from the analysis of baseline immunologic characteristics.

Informed consent for participation in this research was obtained from patients or their guardians. The human experimentation guidelines of the U.S. Department of Health and Human Services were followed in the conduct of this research. Approval for this human study was obtained from the Institutional Review Board of the Massachusetts General Hospital and the ICDDR,B Research and Ethical Review Committees.

**Confirmation of bacterial strains.** All index cases of cholera were confirmed by culturing stool for *V. cholerae* on taurocholate-tellurite-gelatin agar. After overnight incubation of plates, serological confirmation of suspected vibrio colonies was carried out by slide agglutination (18, 21). Rectal swab specimens from household contacts were collected in Cary-Blair transport medium for subsequent plating on taurocholate-tellurite-gelatin agar and colony identification as described above.

**Measurement of vibriocidal and anti-cholera toxin antibody responses in serum.** Vibriocidal antibody assays were performed with methodology previously described, using guinea pig complement and the homologous serotype of *V. cholerae* O1 or O139 as the target organism (19). The concentrations of complement and bacteria have been separately optimized for determining the vibriocidal antibody responses to *V. cholerae* O1 and *V. cholerae* O139. Serum antibodies specific to the cholera toxin B subunit, of both immunoglobulin G (IgG) and IgA isotypes, were measured using a previously described ganglioside GM<sub>1</sub> capture enzyme-linked immunosorbent assay (25).

**Definition of outcomes in household contacts.** Among household contacts, infection was defined as shedding of *V. cholerae* in the stool and/or a fourfold or greater rise in vibriocidal antibody titer during the 21 days of follow-up. Symptomatic illness was defined as infection associated with any diarrheal symptoms; for contacts with a positive rectal swab culture, diarrheal symptoms had to occur within 3 days of the positive result. With the prompt medical care that was afforded by close follow-up of study participants, none of the household contacts required hospitalization or intravenous hydration. Therefore, diarrheal frequency (measured as the number of stools per 24-h period) was used as a marker of disease severity in this group. Severe symptoms among household contacts were defined as eight or more large-volume watery stools per 24-h period, which represented the top quartile of symptomatic household contacts. Because the frequency of stools was recorded on daily home visits only on study days 2 through 7, individuals whose symptoms began more than 7 days after enrollment were not classified for disease severity.

**Statistical analysis.** Analysis was performed using Stata version 8.0 (Stata Corporation, Inc., College Station, Texas) and R version 1.5.0 (<http://www.R-project.org>) (6). The Student *t* test was used to compare log<sub>10</sub>-transformed vibriocidal antibody responses in the index patients. Characteristics of the infected and uninfected household contacts were compared using generalized estimating equations, with an exchangeable correlation matrix, and the reported odds ratios (OR) and *P* values were adjusted for clustering based on household (29). Multivariate analysis was performed using a clustered logistic regression model using the generalized estimating equations, with the final model selected based on backwards elimination. Means are reported in the text and tables with their standard deviation. All reported *P* values are two tailed.

#### RESULTS

**Characteristics of the study population.** Three hundred index patients and 912 of their household contacts were enrolled in the study between January 2001 and September 2004. Of these, 269 patients and 793 contacts completed 21 days of follow-up and were included in this analysis.

The demographic, microbiologic, clinical, and immunologic features of the index cases and household contacts are shown in Table 1. Fifty-eight percent of the index patients with cholera were female, and the mean age was 25 years. Fifty-seven (21%) of the index patients were infected with *V. cholerae* O139, while 212 (79%) were infected with *V. cholerae* O1 El Tor (83 were infected with serotype Ogawa and 129 with serotype Inaba). Forty-three percent of index patients were blood group O; this was significantly higher than the prevalence of blood group O (33%) in the household contacts (*P* = 0.005), despite the familial relatedness. Additionally, the prev-

TABLE 1. Demographic, microbiologic, clinical, and immunologic features of cholera patients and their household contacts

| Characteristic  | Index patients<br>( <i>N</i> = 269) | Household contacts                         |                                   | <i>P</i> (for infected versus<br>uninfected household<br>contacts) |
|---|-------------------------------------|--|-----------------------------------|--|
|   |                                     | Infected <sup>b</sup><br>( <i>N</i> = 260) | Not infected<br>( <i>N</i> = 533) |  |
| Demographic data                                      |                                     |  |                                   |  |
| Age   | 25 ( $\pm$ 14)                      | 18 ( $\pm$ 15)                             | 21 ( $\pm$ 15)                    | 0.03   |
| Female  | 155 (58%)                           | 132 (51%)                                  | 267 (50%)                         | 0.8  |
| Blood group   |                                     |  |                                   |  |
| O   | 115 (43%)                           | 74 (28%)                                   | 190 (36%)                         | 0.03   |
| A   | 60 (22%)                            | 67 (26%)                                   | 136 (26%)                         | 0.9  |
| B   | 72 (27%)                            | 85 (33%)                                   | 155 (29%)                         | 0.4  |
| AB  | 22 (8%)                             | 34 (13%)                                   | 52 (10%)                          | 0.1  |
| Stool or rectal swab culture result                   |                                     |  |                                   |  |
| Any   | 269 (100%)                          | 185 (72%)                                  | NA                                |  |
| O1 Inaba  | 129 (48%)                           | 83 (32%)                                   | NA                                |  |
| O1 Ogawa  | 83 (31%)                            | 53 (20%)                                   | NA                                |  |
| O139  | 57 (21%)                            | 49 (19%)                                   | NA                                |  |
| Severe dehydration                                    | 241 (90%)                           | 0 (0%)                                     | NA                                |  |
| Immunologic data <sup>a</sup>                         |                                     |  |                                   |  |
| Baseline log <sub>10</sub> vibriocidal antibody titer | 1.8 ( $\pm$ 0.7)                    | 1.6 ( $\pm$ 0.9)                           | 1.9 ( $\pm$ 0.9)                  | 0.001  |
| Day 21 log <sub>10</sub> vibriocidal antibody titer   | 3.3 ( $\pm$ 0.6)                    | 2.6 ( $\pm$ 0.9)                           | 1.9 ( $\pm$ 0.9)                  | <0.001   |
| Baseline log <sub>10</sub> anti-CT IgG                | 2.0 ( $\pm$ 0.3)                    | 1.9 ( $\pm$ 0.3)                           | 2.0 ( $\pm$ 0.4)                  | 0.5  |
| Day 21 log <sub>10</sub> anti-CT IgG                  | 2.4 ( $\pm$ 2.4)                    | 2.2 ( $\pm$ 0.3)                           | 2.0 ( $\pm$ 0.3)                  | <0.001   |
| Fourfold rise in vibriocidal antibody titer           | 265 (99%)                           | 190 (73%)                                  | NA                                |  |

<sup>a</sup> The baseline (day 2) immunologic features presented for household contacts exclude those individuals who were symptomatic or had a positive rectal swab for *V. cholerae* upon enrollment in the study (*n* included = 593).

<sup>b</sup> Infection was defined microbiologically (72% of infected contacts), or in the absence of a positive culture, infection was defined serologically by a fourfold increase in vibriocidal antibody titer (28%). NA, not applicable.

absence of blood group O among index cases in our study was significantly higher than that of a randomly selected cohort of individuals from a neighborhood of Dhaka in which cholera is highly endemic who were enrolled in a recent cholera vaccine study at the ICDDR,B (43% in our cohort versus 29% [89 were blood group O out of 310 subjects] in the vaccine study,  $P < 0.001$ ) (20).

The patients admitted to the ICDDR,B and enrolled as index patients in this study had uniformly severe disease regardless of blood group. More than 90% of patients had severe dehydration and received intravenous fluid and antibiotics, and >99% had a fourfold increase in the vibriocidal titer over the 21 days of follow-up. Figure 1 shows the magnitude of the vibriocidal response among index cases, stratified by blood group. Index patients with blood group O had significantly higher vibriocidal titers on day 21 ( $P = 0.02$ ) after the onset of illness than those with other blood groups. Furthermore, in index cases with a baseline vibriocidal antibody titer of  $\leq 80$ , individuals with blood group O had a higher median increase (*n*-fold) in vibriocidal titer than non-blood group O individuals (median increase, 128- versus 64-fold, respectively,  $P = 0.01$ ).

As shown in Table 1, both infected and uninfected household contacts were younger than index patients. The majority of infected household contacts (72%) were identified by the isolation of *V. cholerae* from rectal swab culture; a smaller proportion (28%) was identified by a fourfold increase in the vibriocidal titer without a positive rectal swab culture. Compared to uninfected household contacts, those who developed *V. cholerae* infection were younger, had lower baseline vibrio-

cidal titers, and were less likely to be blood group O. With the prompt provision of medical care by the visiting field team, none of the infected household contacts required hospitalization or intravenous hydration. However, of the 260 infected household contacts, 150 (58%) developed watery diarrhea.

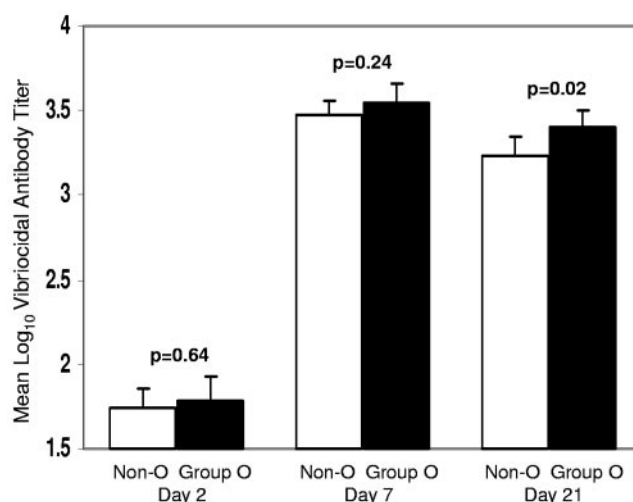


FIG. 1. Vibriocidal antibody responses in index patients with cholera by blood group. Bars represent the mean log<sub>10</sub>-transformed vibriocidal titers, and the error bars represent the 95% CI of the mean. *P* values refer to the differences in mean titer between blood group O and non-blood group O patients on days 2, 7, and 21.

TABLE 2. Risk of infection and severe diarrhea if infected in a cohort of household contacts of index patients with *Vibrio cholerae* O1 and O139 infection

| <i>V. cholerae</i> serogroup in index patient | Blood group of household contact | Infection in household contact <sup>a</sup> | OR for infection (95% CI) | <i>P</i> | Severe diarrhea in household contact, if infected <sup>b</sup> | OR for severe diarrhea (95% CI) | <i>P</i> |
|---|----------------------------------|---|---------------------------|----------|--|---------------------------------|----------|
| O1 & O139 combined                            | O<br>Non-O                       | 74/264 (28%)<br>186/529 (35%)               | 0.80 (0.65–0.98)          | 0.03     | 14/66 (21%)<br>15/173 (9%)                                     | 2.4 (1.2–4.7)                   | 0.01     |
| O1  | O<br>Non-O                       | 49/208 (24%)<br>151/423 (36%)               | 0.67 (0.53–0.85)          | 0.0008   | 8/43 (19%)<br>11/139 (8%)                                      | 2.3 (0.98–5.6)                  | 0.055    |
| O139  | O<br>Non-O                       | 25/56 (45%)<br>35/106 (33%)                 | 1.2 (0.80–1.9)            | 0.32     | 6/23 (26%)<br>4/34 (12%)                                       | 2.3 (0.86–6.2)                  | 0.10     |

<sup>a</sup> Number of household contacts infected/number exposed.<sup>b</sup> Number with severe diarrhea/number infected. Severe diarrhea was defined as eight or more large-volume watery stools per 24 hours. Infected patients were excluded if the onset of symptoms was 8 or more days after enrollment.

The mean number of diarrheal stools among symptomatic household contacts was  $5.5 \pm 4.2$  per day.

**Relationship between blood group O and risk of *V. cholerae* infection among household contacts.** Table 2 shows additional data on the relationship between blood group and the risk of *V. cholerae* infection in our cohort of household contacts of cholera patients. Household contacts with blood group O were less likely to be infected with *V. cholerae* than individuals with other blood groups (OR, 0.80; 95% confidence interval [CI], 0.65 to 0.98;  $P = 0.03$ ). However, once infected, severe symptoms were more likely to occur in household contacts with blood group O (OR, 2.4; 95% CI, 1.2 to 4.7;  $P = 0.01$ ). The mean number of stools in symptomatic contacts with blood group O was  $7.8 \pm 6.5$  per day compared with  $5.1 \pm 4.0$  per day among those with other blood groups ( $P = 0.01$ ).

Table 2 also shows the association between blood group O and the risk of infection stratified by the serogroup of *V. cholerae* to which the contacts were exposed. We found that the association between blood group O and protection from infection was serogroup dependent. While blood group O was associated with protection from infection with *V. cholerae* O1 (OR, 0.67; 95% CI, 0.53 to 0.85;  $P = 0.0008$ ), there was no such protection from infection on exposure to *V. cholerae* O139. However, the associations between blood group O and an increased risk of developing severe diarrheal symptoms when infected were similar for both serogroups of *V. cholerae*. Stratification of the above *V. cholerae* O1 analyses by infecting serotype (Inaba or Ogawa) in contacts of patients with *V. cholerae* O1 showed no significant effect (data not shown).

**Multivariate analysis.** To assess whether the association between blood group O and protection from infection among household contacts was confounded by other factors, such as preexisting adaptive immunity to *V. cholerae*, we performed a stepwise, clustered, multivariate analysis that included baseline vibriocidal antibody titer, baseline anti-cholera toxin antibody titer, blood group, age, and gender. Gender and antitoxin antibody levels were not predictive of the risk of infection and thus were removed by backwards elimination from the final model. Our final model demonstrated that the baseline vibriocidal antibody titer and blood group O were independent predictors of protection from infection with *V. cholerae* O1. Although in the unadjusted analysis, increasing age was associated with protection from infection (Table 1), age was not a

significant predictor after adjustment for other variables in the final model. For each  $\log_{10}$  increase in the baseline vibriocidal antibody titer, there was a 27% decrease in the odds of infection with *V. cholerae* O1 (OR, 0.73; 95% CI, 0.59 to 0.90;  $P = 0.003$ ). Among individuals with blood group O, the odds of infection were half that of non-blood group O individuals in the multivariate model, independent of vibriocidal titer (OR, 0.50; 95% CI, 0.35 to 0.71;  $P = 0.0001$ ). Using the same multivariate model, no significant predictors were found for the risk of infection with *V. cholerae* O139 (data not shown).

## DISCUSSION

The ABH histo-blood group antigens are a set of cellular and secreted glycolipids and glycopeptides that are abundant and widely distributed throughout the body. The ABO phenotype is determined genetically by activity of specific glycosyltransferases and has been shown to be an important determinant of human susceptibility to a number of important gastrointestinal pathogens, including norovirus (14), *Helicobacter pylori* (2), and *V. cholerae*.

Previous studies have demonstrated that blood group O is associated with an increased risk of severe cholera due to the O1 serogroup of *V. cholerae*. In our present cohort, we observed an increased severity of cholera among patients with blood group O infected with both the O1 and O139 serogroups of *V. cholerae*. Additionally, we made the new observation that individuals in Bangladesh with blood group O are protected from infection with *V. cholerae* O1, despite the increased severity of disease once infected. This finding may provide insight into the mechanism of the association between histo-blood group antigens and susceptibility to cholera.

Current models of cholera pathogenesis include two key steps: colonization of the intestine by the organism and the elaboration of cholera toxin (CT), which causes secretory diarrhea. Because the risk of infection with *V. cholerae* O1 is lower in individuals with blood group O, the increased severity of disease once these individuals are infected is unlikely to reflect an increased susceptibility to colonization, as has previously been hypothesized (5). Rather, the increased severity of disease likely reflects a greater susceptibility to later stages of disease pathogenesis, such as an enhanced susceptibility to the secretory effects of CT.



In vitro studies have suggested a biologic mechanism that is consistent with our epidemiologic data, namely, that the A and B histo-blood group glycoconjugates present in non-blood group O individuals may interfere with the binding of CT to its intestinal receptor, the ganglioside GM<sub>1</sub> (15). If this model is correct, then individuals of blood group O should be more susceptible to the pathological effects of CT after colonization, and increased disease severity would be seen in patients infected with either *V. cholerae* O1 or O139, as we have observed.

This interaction between the histo-blood group glycoconjugates and CT is distinct from the interaction with the heat-labile enterotoxin (LT) of enterotoxigenic *Escherichia coli*, despite the homology of the two toxins. While both CT and LT bind to the GM<sub>1</sub> receptor, LT is additionally capable of utilizing the blood group A antigen as an alternate receptor through which the activation of cyclic AMP can be induced (8). This in vitro observation may explain the fact that individuals with blood group O are more susceptible than non-blood group O individuals to severe cholera but not to enterotoxigenic *Escherichia coli*-associated diarrhea (27) as well as provide a potential reason for the evolutionary divergence of the two enterotoxins (28).

Notably, the biological interaction between histo-blood group antigens and CT does not explain why individuals with blood group O are preferentially protected from infection with *V. cholerae* O1 rather than *V. cholerae* O139. Adaptive immunity may account for this observation. In particular, it is possible that the greater severity of *V. cholerae* infection in patients with blood group O may lead to more-potent adaptive immune responses after initial infection and hence enhanced protection from intestinal colonization on subsequent exposure. In support of this, we found that index patients with blood group O in our cohort had significantly higher vibriocidal antibody titers at day 21 following infection.

The hypothesis that adaptive immune responses from prior infection mediate the protection from *V. cholerae* infection conferred by blood group O may also explain our failure to observe an association between blood group O and protection from *V. cholerae* O139 infection. Immunity to *V. cholerae* is serogroup specific, and although substantial preexisting immunity to *V. cholerae* O1 exists in the population in Bangladesh, there is less preexisting immunity to the more recently emerged *V. cholerae* O139 (1). A similar circumstance existed in the early 1980s when the El Tor biotype of *V. cholerae* O1 first appeared and was replacing the classical biotype as the predominant cause of cholera in Bangladesh. The absence of population-level immunity to the El Tor biotype during that time period may account for the failure to detect an association between blood group O and protection from infection in the study from the early 1980s by Glass et al. (10), compared to our present results, in which protection was seen. Similarly, this may explain why no protection from infection was seen in subjects with blood group O when *V. cholerae* O1 El Tor was first introduced into Peru in the 1990s (26).

The prevalence of blood group O among our index cholera patients is lower than that described in previous studies, which reported prevalences ranging from 57 to 64% among patients hospitalized for infection with *V. cholerae* O1 in Bangladesh (5, 7, 10). This may be explained by the fact that after the introduction of a new strain of *V. cholerae* to which the population

is naive, the proportion of cholera patients with blood group O would be expected to decrease over time, as greater adaptive immunity is conferred by previous exposure. Proof of this hypothesis would require further longitudinal studies of a *V. cholerae*-naive population, particularly very young children. A focused analysis of the relationship between blood group and risk of infection with *V. cholerae* among children was not possible in the present study, as there were few young children enrolled in the cohort of household contacts. However, findings reported here do have important implications for the study of infectious disease susceptibility. Specifically, if individuals with an intrinsically increased susceptibility to a disease are also more likely to develop protective adaptive immune responses following exposure, then this enhanced susceptibility may not be identified in the population of an area in which the disease is endemic.

Given our hypothesis regarding the role of adaptive immunity, we anticipated that a multivariate analysis of the risk of cholera infection would demonstrate that the association between blood group O and protection from infection with *V. cholerae* O1 was confounded by the vibriocidal antibody titer on exposure. In fact, our results revealed that blood group O and the baseline vibriocidal antibody titer were independent predictors of protection from *V. cholerae* O1 infection. Although this result may indicate that preexisting adaptive immunity does not explain the relationship between blood group and the risk of *V. cholerae* infection, an alternative explanation is that the vibriocidal antibody titer, which is relatively short-lived after infection, is not the optimal marker of protective immunity following previous infection. Indeed, in a previous analysis of this cohort, we showed that the vibriocidal antibody was an incomplete marker of protection from *V. cholerae* O1 infection (24). We hypothesize that the vibriocidal antibody response may be a surrogate marker for other immune responses that develop following infection and that actually confer protection on subsequent exposure and that these other immune responses are longer lasting than the measured serum vibriocidal antibody titer. Further definition of the specific mechanisms of immunity to *V. cholerae* may allow an improved understanding of the association between blood group and the risk of cholera infection and disease.

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